

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: March 3, 2005, 12:05:14 ; Search time 1226.94 Seconds
(without alignments)
10910.664 Million cell updates/sec

Title: US-09-310-638-1

Perfect score: 2256
Sequence: 1 CCTCTCTCCCTCATCTTTG.....AAAAAAAAAAAAAAAAAAAA 2256

Scoring table: IDENTITY NUC
Gapop 10.0, Gapext 1.0

Searched: 5401638 seqs, 2966923429 residues

Total number of hits satisfying chosen parameters: 10803276

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%
Listing first 45 summaries

Database :

Published Applications NA.*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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3	1719.6	76.2	14446	9	US-09-810-861B-4
4	740	32.8	740	16	US-10-029-386-22811
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6	501	22.2	501	16	US-10-029-386-11696
7	421.4	18.7	2416	16	US-10-032-233-15
8	421.4	18.7	2416	16	US-10-032-233-19
9	421.4	18.7	2416	16	US-10-032-233-33
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11	421.4	18.7	2416	18	US-10-413-432-19

12	421.4	18.7	2416	18	US-10-413-432-33	Sequence 33, Appl1
13	421.4	18.7	2416	18	US-10-324-466-15	Sequence 15, Appl1
14	421.4	18.7	2416	18	US-10-324-466-19	Sequence 19, Appl1
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19	419.8	18.6	2416	9	US-09-748-739A-3	Sequence 3, Appl1
20	419.8	18.6	2416	16	US-10-032-233-11	Sequence 11, Appl1
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22	419.8	18.6	2416	16	US-10-032-233-27	Sequence 27, Appl1
23	419.8	18.6	2416	16	US-10-032-233-35	Sequence 35, Appl1
24	419.8	18.6	2416	18	US-10-413-432-13	Sequence 13, Appl1
25	419.8	18.6	2416	18	US-10-413-432-17	Sequence 17, Appl1
26	419.8	18.6	2416	18	US-10-413-432-27	Sequence 27, Appl1
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31	419.8	18.6	2416	18	US-10-324-466-35	Sequence 35, Appl1
32	418.2	18.5	2381	9	US-09-880-107-2271	Sequence 5, Appl1
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40	418.2	18.5	2416	16	US-10-032-233-39	Sequence 39, Appl1
41	418.2	18.5	2416	16	US-10-032-233-43	Sequence 43, Appl1
42	418.2	18.5	2416	18	US-10-413-432-13	Sequence 13, Appl1
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45	418.2	18.5	2416	18	US-10-413-432-37	Sequence 37, Appl1

ALIGNMENTS

RESULT 1
US-09-810-861B-5
Sequence 5, Application US/09810861B
Patent No. US20020162140A1
GENERAL INFORMATION:
APPLICANT: Mor, Tsafir S.
APPLICANT: Soreq, Hermona
APPLICANT: Amltzen, Charles J.
TITLE OF INVENTION: EXPRESSION OF RECOMBINANT HUMAN ACETYLCHOLINESTERASE IN
FILE REFERENCE: BTI-45
CURRENT APPLICATION NUMBER: US/09/810, 861B
CURRENT FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/190,440
PRIOR FILING DATE: 2000-03-17
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn Ver. 3.1
SEQ ID NO 5
LENGTH: 1725
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: human acetylcholinesterase gene optimized for
OTHER INFORMATION: expression in plants
US-09-810-861B-5

Query Match 76.3%; Score 1722; DB 9; Length 1725;
Best local Similarity 100.0%; Pred. No. 0;
Matches 1722; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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OM nucleic - nucleic search, using sw model

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Searched: 1202784 seqs, 818138359 residues

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-Processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents NA: *
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5: /cgn2_6/prodata/1/ina/PCTUS COMB. seq: *
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SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
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5	2023.2	89.7	2158	4 US-09-949-016-1193	Sequence 1193, Ap
6	1888	83.7	3096	2 US-08-318-826A-6	Sequence 6, Appli
7	1888	83.7	3096	2 US-08-370-156-3	Sequence 3, Appli
8	1888	83.5	3096	2 US-08-814-095-3	Sequence 3, Appli
9	1883.8	83.5	3016	2 US-08-318-826A-7	Sequence 7, Appli
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11	1883.8	83.5	3016	2 US-08-814-095-5	Sequence 5, Appli
12	1845	81.8	1845	3 US-07-732-962A-1	Sequence 1, Appli
13	1845	81.8	1845	5 PCT-US92-06106-1	Sequence 1, Appli
14	1722	76.3	1725	4 US-09-810-861B-5	Sequence 5, Appli
15	1719.6	76.2	5767	4 US-09-810-861B-3	Sequence 3, Appli
16	1719.6	76.2	14446	4 US-09-810-861B-4	Sequence 4, Appli
17	1719.6	76.2	9885	4 US-09-949-016-12934	Sequence 12934, A
18	1357	60.2	9885	4 US-09-949-016-12935	Sequence 12935, A
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20	418.2	18.5	2281	2 US-08-318-826A-9	Sequence 9, Appli
21	418.2	18.5	2416	2 US-08-318-826A-8	Sequence 8, Appli
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27	376	16.7	1215	2 US-08-370-156-26	Sequence 26, Appli

28	374	16.6	374	2 US-08-370-156-24	Sequence 24, Appli
29	371.4	16.5	6867	4 US-09-949-016-17017	Sequence 17017, A
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32	185.6	8.2	764	6 5215909-7	Patent No. 5215909
33	185.6	8.2	764	6 5215909-7	Patent No. 5215909
34	142.6	6.3	3018	1 US-08-347-718B-3	Sequence 3, Appli
35	142.6	6.3	3018	1 US-08-482-262-3	Sequence 3, Appli
36	141	6.2	2734	4 US-09-569-611C-5	Sequence 5, Appli
37	141	6.2	2781	4 US-09-569-611C-6	Sequence 6, Appli
38	141	6.2	3018	6 5200183-1	Patent No. 5200183
39	141	6.2	3018	6 5200183-1	Patent No. 5200183
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45	139.4	6.2	2375	4 US-09-949-016-1976	Sequence 3976, Ap

ALIGNMENTS

RESULT 1
US-08-318-826A-5
Sequence 5, Application US/08318826A

Patent No. 5891725
GENERAL INFORMATION:
APPLICANT: Soreq, Hermona

APPLICANT: Zakut, Haim

APPLICANT: Eckstein, Fritz

TITLE OF INVENTION: Synthetic Antisense
TITLE OF INVENTION: Oligodeoxynucleotides and Pharmaceutical Compositions

NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:

ADDRESSER: Kohn & Associates
STREET: 30500 No. 5891725thwestern Hwy., Suite 410

CITY: Farmington Hills

STATE: Michigan

COUNTRY: US

ZIP: 48334

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/318,826A

FILING DATE:

CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:

NAME: Kohn, Kenneth I.

REGISTRATION NUMBER: 30,955

REFERENCE/DOCKET NUMBER: 2391.00001

TELECOMMUNICATION INFORMATION:

TELEPHONE: (248) 539-5050

TELEFAX: (248) 539-5055

INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 2256 base pairs

TYPE: nucleic acid

STRANDEDNESS: double

TOPOLOGY: linear

MOLECULE TYPE: CDNA to mRNA

HYPOTHETICAL: NO

ANTI-SENSE: NO

ORIGINAL SOURCE:

ORGANISM: Homo sapiens

FEATURE:

OTHER INFORMATION: /note="Splice variant: Exons 1, 2,

OTHER INFORMATION: 3, 4 and 6"

US-08-318-826A-5

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OM nucleic - nucleic search, using sw model

Run on: March 3, 2005, 12:05:14 ; Search time 1683.78 Seconds
(without alignments)
10910.664 Million cell updates/sec

Title:	US-09-310-638-3
Perfect score:	3096
Sequence:	1 CCTCTCTCCCTCATCTTG.....AAAAAAAAAAAAAAAAAAAA 3096

Scoring table: IDENTITY_NUC

Searched: 5401638 seqs, 2966523429 residues
Total number of hits satisfying chosen parameters: 10803276

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Post-processing: Minimum Match 0%

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Database : Published_Applications_NA:*

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SUMMARIES

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2	1720	55.6	5767	9	US-09-810-861B-3	Sequence 3, Appl1
3	1720	55.6	14446	9	US-09-810-861B-4	Sequence 4, Appl1
4	740	23.9	740	16	US-10-029-386-22811	Sequence 22811, A
5	594	19.2	594	16	US-10-029-386-25399	Sequence 25399, A
6	501	16.2	501	16	US-10-029-386-11696	Sequence 11696, A
7	411.2	13.3	449	10	US-09-918-995-27556	Sequence 27556, A
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11	383.2	12.4	2416	18	US-10-413-433-15	Sequence 15, Appl

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39	380	12.3	2416	16	US-10-032-233-31	Sequence 31, Appl
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35	380	12.3	2416	9	US-09-748-739A-7	Sequence 7, Appl
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33	380	12.3	2381	9	US-09-880-107-2271	Sequence 2271, Ap
32	381.6	12.3	2416	18	US-10-324-466-35	Sequence 35, Appl
31	381.6	12.3	2416	18	US-10-324-466-27	Sequence 27, Appl
30	381.6	12.3	2416	18	US-10-324-466-11	Sequence 11, Appl
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23	381.6	12.3	2416	16	US-10-032-233-17	Sequence 17, Appl
22	381.6	12.3	2416	16	US-10-032-233-17	Sequence 17, Appl
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20	381.6	12.3	2416	9	US-09-748-739A-3	Sequence 3, Appl
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16	383.2	12.4	2416	18	US-10-324-466-13	Sequence 13, Appl
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ALIGNMENTS

RESULT 1

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? Patent No. US20020162140A1
? GENERAL INFORMATION:
? APPLICANT: Mor, Tsafir S.
? APPLICANT: Soreq, Hermona
? APPLICANT: Artztzem, Charles J.
? APPLICANT: Mason, Hugh S.
? TITLE OF INVENTION: EXPRESSION OF RECOMBINANT HUMAN ACETYLCHOLINESTERASE IN
? TITLE OF INVENTION: TRANSGENIC PLANTS
? FILE REFERENCE: BTI-45
? CURRENT APPLICATION NUMBER: US/09/810,861B
? CURRENT FILING DATE: 2001-03-16
? PRIOR APPLICATION NUMBER: 60/190,440
? PRIOR FILING DATE: 2000-03-17
? NUMBER OF SEQ ID NOS: 5
? SOFTWARE: PatentIn Ver. 3.1
? SEQ ID NO 5
? LENGTH: 1725
? TYPE: DNA
? ORGANISM: Artificial Sequence
? FEATURE:
? OTHER INFORMATION: Description of Artificial Sequence: synthetic
? OTHER INFORMATION: human acetylcholinesterase gene optimized for
? US-09-810-861B-5

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Query Match	Score	DB	Length
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Matches 1723; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Issued Patents NA: *
1: /cgn2_6/ptodata/1/ina/5A.COMB.seq: *
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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9	1888	61.0	2256	2	US-08-814-095-1
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13	1729	55.8	1845	5	PCT-US92-06106-1
14	1722.4	55.6	1725	4	US-09-810-861B-5
15	1720	55.6	5767	4	US-09-810-861B-3
16	1720	55.6	14446	4	US-09-810-861B-4
17	1529.6	49.4	9885	4	US-09-949-016-12934
18	1529.6	49.4	9885	4	US-09-949-016-12935
19	1528	49.4	35060	3	US-08-814-095-7
20	1215	39.2	1215	2	US-08-370-156-25
21	380	12.3	2381	2	US-08-318-826A-9
22	380	12.3	2416	2	US-08-318-826A-8
23	380	12.3	2416	3	US-09-334-489-1
24	380	12.3	2416	3	US-09-334-489-2
25	380	12.3	2444	4	US-09-949-016-5275
26	378.4	12.2	2400	6	5215909-13
27	378.4	12.2	2400	6	5215909-13

28	374	12.1	374	2	US-08-370-156-24	Sequence 24, Appl
29	371.4	12.0	68667	4	US-09-949-016-17017	Sequence 17017, A
30	322.8	10.4	2445	6	5215909-9	Patent No. 5215909
31	322.8	10.4	2445	6	5215909-9	Patent No. 5215909
32	185.6	6.0	764	6	5215909-7	Patent No. 5215909
33	185.6	6.0	764	6	5215909-7	Patent No. 5215909
34	142.6	4.6	3018	1	US-08-347-718B-3	Sequence 3, Appl1
35	142.6	4.6	3018	1	US-08-482-262-3	Sequence 3, Appl1
36	141	4.6	2734	4	US-09-569-611C-5	Sequence 5, Appl1
37	141	4.6	2781	4	US-09-569-611C-6	Sequence 6, Appl1
38	141	4.6	3018	6	5200183-1	Patent No. 5200183
39	141	4.6	3018	6	5200183-1	Patent No. 5200183
40	139.4	4.5	1746	4	US-10-023-515-3	Sequence 3, Appl1
41	139.4	4.5	2158	4	US-10-023-515-1	Sequence 1, Appl1
42	139.4	4.5	2184	1	US-08-445-050-8	Sequence 8, Appl1
43	139.4	4.5	2184	1	US-08-204-691-8	Sequence 8, Appl1
44	139.4	4.5	2344	3	US-09-347-878-31	Sequence 31, Appl1
45	139.4	4.5	2375	4	US-09-949-016-3976	Sequence 3976, Ap

ALIGNMENTS

RESULT 1
US-08-318-826A-6
Sequence 6, Application US/08318826A
Patent No. 5891725
GENERAL INFORMATION:
APPLICANT: Soreq, Hermona
APPLICANT: Zakut, Haim
APPLICANT: Eckstein, Filtz
TITLE OF INVENTION: Synthetic Antisense
TITLE OF INVENTION: Oligodeoxynucleotides and Pharmaceutical Compositions
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kohn & Associates
STREET: 30500 No. 589175thwegtein Hwy., Suite 410
CITY: Farmington Hills
STATE: Michigan
COUNTRY: US
ZIP: 48334
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,826A
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Kohn, Kenneth I.
REGISTRATION NUMBER: 30,955
REFERENCE/DOCKET NUMBER: 2391.00001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (248) 539-5050
TELEFAX: (248) 539-5055
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 3096 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: 160..1959
OTHER INFORMATION: /note=Splice variant: Exons 1, 2,

GenCore version 5.1.6
Copyright (c) 1993 - 2005 CompuGen Ltd.

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Run on: March 3, 2005, 06:55:38 ; Search time 487,289 Seconds
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Title: US-09-310-638-5

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Gapop 10.0 , Gapext 1.0

Searched: 1202784 segs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
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Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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8	1883.8	62.5	2256	2	US-08-370-156-1
9	1883.8	62.5	2256	3	US-08-814-095-1
10	1748	58.0	2158	4	US-09-949-016-1192
11	1748	58.0	2158	4	US-09-949-016-1193
12	1724.8	57.2	1845	1	US-07-732-862A-1
13	1724.8	57.2	1845	5	PCT-US82-06106-1
14	1722	57.1	1725	4	US-09-810-861B-5
15	1720.2	57.0	1446	4	US-09-810-861B-4
16	1720.2	57.0	1446	4	US-09-810-861B-3
17	1359.6	45.1	9885	4	US-09-849-016-12934
18	1359.6	45.1	9885	4	US-09-849-016-12935
19	1358	45.0	35060	3	US-08-814-095-7
20	1135.8	37.7	1215	2	US-08-370-156-26
21	380	12.6	2381	2	US-08-318-826A-9
22	380	12.6	2416	2	US-08-318-826A-8
23	380	12.6	2416	3	US-09-334-489-1
24	380	12.6	2416	3	US-09-334-489-2
25	378.4	12.5	2400	6	US-09-949-016-5275
26	378.4	12.5	2400	6	US-09-949-016-5275
27	378.4	12.5	2400	6	US-09-949-016-5275

28	374	12.4	374	2	US-08-370-156-24	Sequence 24, Appl
29	371.4	12.3	6867	4	US-09-949-016-17017	Sequence 17017, A
30	322.8	10.7	2445	6	5215909-9	Patent No. 5215909
31	322.8	10.7	2445	6	5215909-9	Patent No. 5215909
32	185.6	6.2	764	6	5215909-7	Patent No. 5215909
33	185.6	6.2	764	6	5215909-7	Patent No. 5215909
34	144.2	4.8	3018	1	US-08-347-718B-3	Sequence 3, Appl1
35	144.2	4.8	3018	1	US-08-482-262-3	Sequence 3, Appl1
36	142.6	4.7	2734	4	US-09-589-611C-5	Sequence 5, Appl1
37	142.6	4.7	2781	4	US-09-589-611C-6	Sequence 6, Appl1
38	142.6	4.7	3018	6	5200183-1	Patent No. 5200183
39	142.6	4.7	3018	6	5200183-1	Patent No. 5200183
40	141	4.7	2184	1	US-08-445-050-8	Sequence 8, Appl1
41	141	4.7	2184	1	US-08-204-691-8	Sequence 31, Appl1
42	141	4.7	2344	3	US-09-347-878-31	Sequence 31, Appl1
43	141	4.7	2375	4	US-09-949-016-3976	Sequence 3976, Ap
44	141	4.7	2428	1	US-08-445-050-1	Sequence 1, Appl1
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ALIGNMENTS

RESULT 1
US-08-318-826A-7
Sequence 7, Application US/08318826A
Patent No. 5891725
GENERAL INFORMATION:
APPLICANT: Soreq, Hermona.
APPLICANT: Zakut, Haim
APPLICANT: Eckstein, Fritz
TITLE OF INVENTION: Synthetic Antisense
TITLE OF INVENTION: Oligodeoxynucleotides and Pharmaceutical Compositions
TITLE OF INVENTION: Containing Them
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSER: Kohn & Associates
STREET: 30500 No. 5891725thwestern Hwy., Suite 410
CITY: Farmington Hills
STATE: Michigan
COUNTRY: US
ZIP: 48334
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/318,826A
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Kohn, Kenneth I.
REGISTRATION NUMBER: 30,955
REFERENCE/DOCKET NUMBER: 2391.00001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (248) 539-5050
TELEFAX: (248) 539-5055
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 3016 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: 160..2010
OTHER INFORMATION: /note="Splice Variant: Exons 1, 2,

GenCore version 5.1.6
Copyright (c) 1993 - 2005 CompuGen Ltd.

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Title: US-09-310-638-5

Perfect score: 3016

Sequence: 1 CCTCTCTCCCTCATCTTTG.....AAAAAAAAAAAAAAAAAAAA 3016

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Total number of hits satisfying chosen parameters: 10803276

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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C	594	19.7	594	US-10-029-386-25399	Sequence 25399, A
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24	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
25	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
26	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
27	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
28	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
29	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
30	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
31	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
32	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
33	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
34	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
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36	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
37	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
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39	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
40	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
41	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
42	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
43	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
44	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1
45	383.2	12.7	2416	US-10-324-466-15	Sequence 15, Appl1

ALIGNMENTS

RESULT 1

US-09-810-861B-5

Sequence 5, Appl1 US/09810861B

Patent No. US20020162140A1

GENERAL INFORMATION:

APPLICANT: Mor, Tsafir S.

APPLICANT: Soreq, Hermona

APPLICANT: Amlitz, Charles J.

APPLICANT: Maon, Hugh S.

TITLE OF INVENTION: EXPRESSION OF RECOMBINANT HUMAN ACETYLCHOLINESTERASE IN TRANSGENIC PLANTS

FILE REFERENCE: BTI-45

CURRENT APPLICATION NUMBER: US/09/810, 861B

CURRENT FILING DATE: 2001-03-16

PRIOR APPLICATION NUMBER: 60/190,440

PRIOR FILING DATE: 2000-03-17

NUMBER OF SEQ ID NOS: 5

SOFTWARE: PatentIn Ver. 3.1

SEQ ID NO 5

LENGTH: 1725

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURES:

OTHER INFORMATION: Description of Artificial Sequence: synthetic

OTHER INFORMATION: human acetylcholinesterase gene optimized for

OTHER INFORMATION: expression in plants

US-09-810-861B-5

Query Match

Best Local Similarity 100.0%; Pred. No. 0;

Matches 1722; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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